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Perfect score:
                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                  Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                             Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                     262574 seqs, 29422922 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-091-628-2
Issued_Patents_AA:*

1: /ggn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9, 2003, 07:07:25; Search time 19 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .....PGPMDCHRALEPVGHITSCE 377
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583.813 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match I	Length	BB	ID	Description
	884	44.7	348	_	US-08-176-126B-2	Sequence 2, Appli
2	œ	44.7	348	N	-08-669-435-2	e 2
ω	884	4	348	σ	T-US94-14431	equence
4	0	43.5	348	_	8-176-126B-4	Α
v	860.5	43.5	348	N	-08-669-	equence 4,
σ	.0	43.5	348	ហ	T-US94-14431	4
7	101	•	370	4	US-08-405-271A-21	21,
8	0	5.1	424	4	9	5009
9	99.5	5.0	387	4	US-08-993-088A-7	7,
10	S	•	387	4	US-08-993-424B-7	Sequence 7, Appli
11	98.5	5.0	344	ω	US-09-110-116-4	e 4,
12	. 98	•	835	4	US-09-284-819-6	ο,
13	96.5	4.9	3177	N	-451-	4
14	96	4.9	521	4	-	4
15	96	4.9	572	4.	US-08-937-067-13	13
16	96	4.9	652	ω	÷	۳
17	96	4.9	652	4	56-322-	2
18	93	4.7	372	N	US-08-514-451A-9	Sequence 9, Appli
19	93	4.7	372	4	US-09-048-916B-8	е 8
.20	93	4.7	487	_	١,	e 2,
21	93	4.7	532	۳	US-08-140-729A-3	е 3,
22	93	4.7	532	ш	8-546-6	е Э,
23	93	4.7	532	N	-08-916-745-	е Э,
24	93	4.7	532	N	US-09-042-929-3	e u
25	93	4.7	532	N	-661-	e u
26	93	4.7	532	N	-09	Sequence 3, Appli
27	93	4.7	532	ω	US-09-198-650-3	'n

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63 RRPWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVDGD 122

7 SSSACPANSS--EEELPVGLEVHGN--LELVFTVVSTVMMGLLMFSLGCSVEIRKLWSHI 62 NSSICNPNATICEGDSCIAPESNFNAILSVVMSTVLTILLALVMFSMGCNVELHKFLGHL 62

44.7%; Score 884; DB 1; Length 348; 46.9%; Pred. No. 2.4e-82; ative 74; Mismatches 102; Indels

Indels 10;

Gaps

Query Match Best Local Similarity

Matches 164; Conservative

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28
90	90.5	91	91.5	91.5	91.5	91.5	91.5	91.5	91.5	91.5	91.5	91.5	92	92.5	93	93	93
4.5	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.6	4.7	4.7	4.7	4.7
372	405	843	485	485	485	485	372	372	372	367	367	367	683	554	532	532	532
ω	4	4	ហ	N	μ	ب	4	4	4.	w	w	N	4	4	4	w	w
US-09-170-331-6	US-09-134-001C-4999	US-09-361-631-1	PCT-US93-08174-15	US-08-452-930-15	US-08-086-631-15	US-08-453-956-15	US-09-113-426-4	US-09-351-198-4	US-08-188-275A-4	US-08-676-351-3	US-08-454-552-4	US-08-454-549-4	US-09-134-001C-5576	US-09-134-001C-5109	US-09-042-709A-3	US-09-042-937-3	US-09-042-913-3
Sequence 6, Appl	Sequence 4999,	Sequence 1, i	Sequence 15,	Sequence 15,	•	Sequence 15,	Sequence 4, i	Sequence 4, i	Sequence 4, 1	Sequence 3, 1	Sequence 4, 1	Sequence 4, 1	Sequence 5576	Sequence 5109,	Sequence 3, 1	Sequence 3, 1	Sequence 3, 1
Appli	9, Ap	1, Appli	Appl	App1	Appl	Appl	Appli	Appli	Appli	Appli	Appli	Appli	6, Ap	9, Ap	Appli	Appli	appli

ALIGNMENTS

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US-08-176-126B-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence-2, Application_US/08176126B
Patent No. 5589358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                               INFORMATION FOR SEQ ID NO:
                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acid
        TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/176,126B
FILING DATE: 29-DEC-1993
                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBERTUR SURVEY ADDRESS:
CORRESPONDENCE ADDRESS:
Arnold, White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:—Dawson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TUMBER-OF-SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
ZIP: 77210
                                                                                                                                                                                 NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: WAKE:002/PAR
                                                                                                                 TELEPHONE: (512) 474-7577
                                                                                                                                                                                                                                                   FILING DATE: 2:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: ALICE., ADDRESSEE: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Houston
                                             amino acid
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                                                              348 amino acids
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183 123

123 MDLSISMTTCSTVAALGMMPLCIYLYTWSWSLQQNLTIPYQNIGITLVCLTIPVAFGVYV 182

182

122

RRPWGIVVGFLCQFGIMPLTGFVLSVAFGILPVQAVVVLIQGCCPGGTASNILAYWVDGD

NYRWPKQSKIILKIGAVVGGVLLLVVAVAGVVLAKGSWNSDITLLTISFIFPLIGHVTGF 242 MDLSVSMTTCSTLLALGMMPLCLFIYTKMWVDSGTIVIPYDSIGTSLVALVIPVSIGMYV 밁

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                                                                                                                                   TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-669-435-2
                                                                   Matches
                                                                                Query Match
Best Local Similarity
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Natent No. 5869265
GENERAL-INFORMATION:
                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8-669-435-2
                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
LENGTH: 348 amino aci
                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: PAXKET, DAVID I.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: WAI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Dawson, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                  LENGTH: 348 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 29-DEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 26-JU
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                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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                                                                 164;
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                                 7 SSSACPANSS
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NSSI CNPNATI CEGDSCIAPESNENAI LSVVMSTVLTILLALVMESMGCNVELHKELGHL
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P.O. Box 4433
                                                                                                                                                                                                                                                                         (512)
                                                                 Conservative
                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          METHODS
                                                                                 44.78;
                              -EEELPVGLEVHGN--LELVFTVVSTVMMGLLMFSLGCSVEIRKLWSHI
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                                                                                                                                                                                                                                         2
                                                                74;
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                                                                Score 884; DB 2;
Pred. No. 2.4e-82;
4; Mismatches 102
                                                                                                 Length 348;
                                                                 Indels
                                                                10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AND
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62
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SSACPANSS--EEELPVGLEVHGN--LELVFTVVSTVMMGLLMFSLGCSVEIRKLWSHI 62

NSSICNPNATICEGDSCIAPESNFNAILSVVMSTVLTILLALVMFSMGCNVELHKFLGHL

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RESULT 3
PCT-US94-14431A-2
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                                                                     PCT-US94-14431A-2
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Matches
                Query Match
Best Local Similarity
                                                                                                                                                      TELEPAX: (713) 789-2679
TELEX: 79-0924(1) GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Relei
SOFTWARE: #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT.
FILING DATE: 29-DEC-199-
                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 29-DEC-11
CLASSIFICATION:
                                                                                                                                                                                                                                          NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: WAI
TELECOMMUNICATION INFORMATION:
                                                                                 TOPOLOGY: 13
                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION: NAME: PARKER, DAVID L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U
ZIP: 77210
                                                                                                                                                                                                         TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NHKWPQKAKIILKIGSIAGAILIVLIAVVGGILYQSAWTIEPKLWIIGTIYPIAGYGLGF
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                                                                                                                                    348 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arnold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                                   protein
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                                                                                                                                                                                                                                                                                                                                                                                                                       29-DEC-1994
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Box 4433
                44.7%;
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                                                                                                                                                                                                                                                                                                                                                                    USSN 08/176,126
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74;
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              Score 884; DB 5;
Pred. No. 2.4e-82;
 Mismatches
 102;
                                Length 348;
   10;
 Gaps
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RRPWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVDGD

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RRPWGIVVGFLCQFGIMPLTGFVLSVAFGILFVQAVVVLIQGCCPGGTASNILAYWVDGD

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                                                                                                                                                                                                  ; MOLECULE TYPE: protein US-08-176-126B-4
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                                                                                                                             Query Match (1835) Score 860.5; DB 1
Best Local Similarity (1856) Pred. No. 6.2e-80;
Matches 160; Conservative 68; Mismatches 104
                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acid
                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000
                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: WAKE:002/PAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                  TELEPHONE: (512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/0 FILING DATE: 29-DEC-1993
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ZIP: 77210
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PWGICVGFLCQFGIMPLTGFILSVAFDILPLQAVVVLIIGCCPGGTASNILAYWVDGDMD
                  PWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVDGDMD
                                                                   CSGASCVVPESNFNNI --
                                                                                         CSSSSACPANSSEEELPVGLEVHGNLELVFTVVSTVMMGLLMFSLGCSVEIRKLMSHIRR
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                                                               -LSVVLSTVLTILLALVMFSMGCNVEIKKFLGHIKR
                                                                                                                                                              DB 1;
                                                                                                                                 104;
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                                                                                                                                                                Length 348;
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Patent (No. 1586926
GENERAL TANTE ORMAN
APPLICANT: D
                                                                                                                    Matches
                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
SOFTWARE: PatentIn Release #1.0, Vers
CURRENT APPLICATION DATA:
OUR CONTROL WITCH.
                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acids
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                       TELEPHONE: (512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                     NAME: Parker, David L.
REGISTRATION NUMBER: 32,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                      TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 29-DEC-1993
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65 PWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVDGDMD 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VAAYQTYKRRIKNKHGKKNSGCTEVCHTRKSTSSRETNAFLEVNEEGAITP 355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ARIAGLEWYRCRTVAFETGMONTOLCSTIVOLSFTPEELNVVFTFPLIYSIFOLAFAAIF 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ALFTHOSWORCRTISLETGAONIOMCITMLOLSFTABHLVOMLSFPLAYGLFOLIDGFLI 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LSVSMTTCSTLLALGMMPLCLLIYTKMWVDSGSIVIPYDNIGTSLVALVVPVSIGMFVNH 184
                                                                              CSSSSACPANSSEEELPVGLEVHGNLELVFTVVSTVMMGLLMFSLGCSVEIRKLWSHIRR
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P.O. Box 4433
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                                                                                                                    Conservative
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ILEAL BILE ACID TRANSPORTER COMPOSITIONS METHODS
                                                                                                                                  43.5%;
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                                                                                                                                                                                                                                                                                       4.
                                                                                                                 68;
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                                                                                                                 Score 860.5; DB 2;
Pred. No. 6.2e-80;
B; Mismatches 104;
                                                  ----LSVVLSTVLTILLALVMFSMGCNVEIKKFLGHIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Version #1
                                                                                                                   Indels 19;
                                                                                                                                                Length 348;
                                                                                                                 Gaps
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                                                                                                                                                                           PCT-US94-14431A-4
                                                                                                              Matches 160;
                                                                                                                           Query Match
Best Local S
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                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 29-DEC-
                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,1
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 29-DE0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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PWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVDGDMD
                                                                               CSSSSACPANSSEEELPVGLEVHGNLELVFTVVSTVMMGLLMFSLGCSVEIRKLWSHIRR
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                                                 CSGASCVVPESNFNNI---
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                                                                                                                Conservative
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29-DEC-1993
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                                                                                                                           Score 860.5;
Pred. No. 6.
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                                                                                                              Mismatches
                                                 6.2e-80;
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                                                                                                                                            Length 348;
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                                                                      Matches
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                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: EVANS, CHRISTOPHER J.
APPLICANT: KEITH, DUANE E.
TITLE OF INVENTION: OPIOID RECEPTOR GENES
NUMBER OF SEQUENCES: 25
                                                                                                                                                                                                                                                                                                                                      TELEX: 90-4030 MRSNFO:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                         FEATURE:
NAME/KEY: Modified-si
LOCATION: group(18, 3
OTHER INFORMATION: /n
OTHER INFORMATION: th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: MURASHIGE, KATE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 22
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                ELECUTATION (202)
                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MURASHIGE, KATE H. REGISTRATION NUMBER: 29,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                  LENGTH: 370 amino acids TYPE: amino acid STRANDEDNESS: single
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                                                                      91;
                                 3 ANCSSS--SACP---ANSSEEELPVGLEVHGNLELVFTVVSTVMMGLLMFSLGCSVEIRK 57
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                                                                                        Similarity
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                                                                    Score 101; DB 4;
Pred. No. 0.037;
8; Mismatches 147
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for N-linked glycosylation"
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RESULT 9
US-08-993-088A-7
; Sequence 7, Application US/08993088A
; Patent No. 6287855
; Patent No. 6287857
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LENGTH: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/134,001C CURRENT FILING DATE: 1998-08-13 PRIOR APPLICATION NUMBER: US 60/064,964 PRIOR FILING DATE; 1997-11-08 PRIOR APPLICATION NUMBER: US 60/055,779 PRIOR FILING DATE: 1997-08-14 NUMBER OF SEQ ID NOS: 5674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Staphylococcus epidermidis
09-134-001C-5009
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              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: GTC-007
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o. 6380370
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                                                                                                                                                                                                                                                                                                                                        AY-----LLAISFSLKP----VQAIAVLIMGCCPG-GTISNIFTFWV-DGDMDLSISMTT 131
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28.4%; Pred. No. 0.05;
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RESULT 10
US-08-993-424B-7
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OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/993,088A
FILING DATE: 18-DEC-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/033,851
APPLICATION NUMBER: 60/033,851
APPLICATION NUMBER: 30
PRIOR APPLICATION NUMBER: 39,179
REFERENCE/DOCKET NUMBER: 19846
REGISTRATION NUMBER: 19846
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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ZIP: 07065-0900
COMPUTER READABLE FORM:
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APPLICANT: Sullivan, Kathleen
TITLE OF INVENTION: GALANIN RECEPTOR GALR2 AND
TITLE OF INVENTION: NUCLEOTIDES ENCODING SAME
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 732-594-1958
TELEFAX: 732-594-4720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.0%; Score 99.5; DB 4; Length 387; Local Similarity 20.8%; Pred. No. 0.056; see 73; Conservative 47: Mismath.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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            Application US/08993424B
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                                                                                                           G--RVCAAARGTHSGSVLERESSDLLHMSEAAGALRPCPGASQPC--ILEP
                                                                                                                                             GCTEVCHTRKSTSS-----RETNAFLEVNE-EGAITPGPPGPMDCHRALEP 369
                                                                                                                                                                                                                        FPLAYGLFQLIDGFLIVAAYQTYKR-----RLKNKHGKKN--
                                                                                                                                                                                                                                                               GLTYARTLRYLWRAVDPVAAGSGARRAKRKVTRMILIVAALFCLCWMPHHALILCVWFGQ 263
                                                                                                                                                                                                                                                                                                   GFLLALFTHQSWQRCRTISLETGAQNIQMCITMLQLSFTA----EHLVQML----S
                                                                                                                                                                                                                                                                                                                                            IGLIWGLSLLFSGPYLSYYRQSQLANLTVCHPAWSAPRRAMDICTFVFSYLLPVL--VL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTVGNTLVLAVLLRGGQAVS---TINLFILNLGVADLCFIL-----CCVPFQATI 85
                                                                                                                                                                                    FPLTRATYAL----RILSHLVSYANSCVNPIVYALVSKHFRKGFRTICAGLLGRAPGRAS
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P.O. Box 2000, 1
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Best Local :
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NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 1984
TELECOMMUNICATION INFORMATION:
TELEPHONE: 732.594-1958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 387 amino acids
TYPE: amino acid
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CORRESPONDENCE ADDRESS:
ADDRESSEE: March , -
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/033,851
FILING DATE: 27-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
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OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version
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TOPOLOGY: lir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                      325
                                                                             264
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                                                                                                                                                                                                                                    146 İGLIMĞLŞİLFSGPYLSYYROSQLANLTVCHPAWSAPRRRAMDİCTFVFSYLLFVL--VL 203
320
                                                                                                                                                                                           241 GFLLALFTHQSWQRCRTISLETGAQNIQMCITMLQLSFTA-----EHLVQML-----S
                                                                                                                                                                                                                                                                           196 IGAVVGGVLLL-------VVAVAGVVLAKGSWNS-----DITLLTISFIFPLIGHVT 240
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Similarity 20.8%;
73; Conservative 4:
                                      GCTEVCHTRKSTSS----RETNAFLEVNE-EGAITPGPPGPMDCHRALEP 369
                                                                                                                                                         GLTYARTLRYLWRAVDPVAAGSGARRAKRKVTRMILIVAALFCLCWMPHHALILCVWFGQ 263
  G--RVCAAARGTHSGSVLERESSDLLHMSEAAGALRPCPGASQPC--ILEP 366
                                                                             FPLTRATYAL----RILSHLVSYANSCVNPIVYALVSKHFRKGFRTICAGLLGRAPGRAS
                                                                                                                  FPLAYGLFQLIDGFLIVAAYQTYKR-----RLKNKHGKKN---
                                                                                                                                                                                                                                                                                                                                                                                             GTVGNTLVLAVLLRGGQAVS---TTNLFILNLGVADLCFIL------CCVPFQATI 85
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Pred. No. 0.056;
7; Mismatches 118;
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RESULT 11

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APPLICANT: The Government of the United States of America
APPLICANT: as represented by The Secretary of the
APPLICANT: Department of Health and Human Services
ITILE OF INVENTION: Methods and Compositions for Inhibiting Inflammation
ITILE OF INVENTION: and Angiogenesis Comprising a Mammalian CD97 Alpha
ITILE OF INVENTION: Subunit
FILE REFERENCE: 015280-263100US
CURRENT APPLICATION NUMBER: US/09/284,819
CURRENT APPLICATION NUMBER: US/09/284,819
CURRENT FILING DATE: 1999-08-20
EARLIER APPLICATION NUMBER: US 60/027,871
EARLIER APPLICATION NUMBER: W5 60/027,871
EARLIER APPLICATION NUMBER: W5 60/027,871
FEARLIER FILING DATE: 1996-10-25
LARLIER FILING DATE: 1997-10-24
NUMBER OF SEQ ID NOS: 21
SOFTWARE: PatentIn Ver. 2.1
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US-09-284-819-6
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                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6, Application US/09284819 Patent No. 6365712
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Best Local (
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APPLICANT: Cohan, Victoria L.
APPLICANT: Scuart, Susan G.
TITLE OF INVENTION: HUMAN EMR1-LIKE G PROTEIN COUPLED TITLE OF INVENTION: RECEPTOR
FILE REFERENCE: PF-0550 US
CURRENT APPLICATION NUMBER: US/09/110,116
CURRENT FILING DATE: 1998-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kelly, Kathleen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             127 ISMTTCSTVAAL------GMMPLC------IYLYTWSWSL------QQNLT 159
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                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: McClung, Barbara G.
REGISTRATION NUMBER: 33,113
REFERENCE/DOCKET NUMBER: 0335
TELECOMMUNICATION INFORMATION:
TELEPHONE: 510-601-2708
                                                                      TELEFAX: 510-655-3542
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: CD97amino acid sequence encoded by full-length OTHER INFORMATION: clone paT276
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                               SEQUENCE CHARACTERISTICS:
LENGTH: 3177 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                           FILING DATE:
STRANDEDNESS:
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                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                            INFORMATION FOR SEQ ID NO:
                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 521 amino acids
                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MAO, JOYCE
TITLE OF INVENTION: CDNA CLONE HAPOI67 THAT
TITLE OF INVENTION: A HUMAN 7-TRANSMEMBRANE
                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/049,329
FILING DATE: 11-JUN-1997
ATTURNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/956,322
FILING DATE: 23-OCT-1997
CLASSIFICATION: 536
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                                                                                                                                                                            NAME: PRESTIA, PAUL F
REGISTRATION NUMBER: 23
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER:
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                            STRANDEDNESS:
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                                               amino acid
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US-08-956-322-4

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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 23647-20018.00
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-937-067-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13, Application US/08937067 Patent No. 6433155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       ZIP: 94304-1018
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: MORRISON & FOERSTER STREET: 755 Page Mill Road CITY: Palo Alto
                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      352 AITPG 356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  408 SEVSTIQNTRMLAFKATAQLFILGCTWCLGLLQVGPAAQ-----VMAY-LFTIINSL 458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            348 VAISAASWPHLYGTADRCWLHLDQGFMWSFLGPVCAIFSANLVLFILVFWILKRKLSSLN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              213 VVLAKGSW-----NSDITLLTI--SFIFPLIGHVTGFL---LALFTHQSW------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       316 TARNLTVVNYSSINRLMKWIMFPVGYG-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121 GDMDLSISMTTCSTVAALGMM------PLC-----IYLYTWSWSL-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               203 NCSHLSSFAVLMALTSQEEDP------VLTVITYV-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 HIRRPWGIAVGLLCQFGLMPFTAYLLAISFSLKPVQAIAVLIMGCCPGGTISNIFTFWVD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 NC---SSSSACPANSSEEELPVGLEVHGNLELVFTVVSTVMMGLLMFSLGCSVEIRKLWS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QGFF1FLVYCLLSQQVQKQYQKWFREIVKSKSESETYTLSSKMGPDSKPS---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ----QRCRTISLETGAQ----NIQMCITMLQLSFTAEHLVQMLSFPLAYGLFQLID--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -QQNLT-IPYQNIGITLVCLTIPVAFGVYVNYRWPKQSKIILKIGAVVGGVLLLVVAVAG 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ---SLHLQLSLCLFLAHLLFLVGIDRTEPKVLCSIIAGALHYLYLAAFTWMLLEGVHLFL
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                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Melkonyan,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Umansky, Samuil
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A FAMILY OF GENES ENCODING
APOPTOSIS-RELATED PEPTIDES;
METHODS OF USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.9%;
                                                                                                                                                                                                                  US/08/937,067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 96; DB 4; Length 521; Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches 107;
                                                                                                                                                                                                                                                       Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -KNSGCTEVCHTRKSTSSRETNAFLEVNEEG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ----CK--AIQNTST----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PEPTIDES ENCODED THEREBY AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels 184;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -VPAVT 347
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                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear ; MOLECULE TYPE: peptide US-08-937-067-13
                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
 538
                                                                                                                          433
                              237
                                                                                                                                                                                      391 TITILAMG-----
                                                             480 AT--IVLACYFYEQAFREHWERTWILLQTCKSYAVPCPPRHFSPMSPDFTVFMIKYLMTMI
                                                                                           209 AVAGVVLA-----
                                                                                                                                                     154 LQQNLTIPYQNIGITLVCLTIPVAFGV-----YVNYRWPKQSKIILKIGAVVGGVLLLVV 208
                                                                                                                                                                                                                                                  337 LFMVLYFFGMASSI----WWVILSLTWFLAAGM--KWGHEAIEANSQYFHLAAWAVPAVK 390
                                                                                                                                                                                                                  97 AIAVLIMGCCPGGTISNIFTFWVDGDMDLSISMTTCSTVAALG---MMPLCIYLYTWSWS 153
                                                                                                                                                                                                                                                                                                               57;
                                                                                                                                                                                                                                                                                40 VMMGLLMFSLGCSVEIRKLWSHIRRPWGIAVGLLCQFGLMPFTA---YLLAISFSLKPVQ 96
VGITTGFWIWSGKTLQSWRR 557
                             -GHVTGF-LLALFTHQSWQR 254
                                                                                                                        -----IGTSFLLAGFVSLFRIRTIMKHDGTKTEKLEKLMVRIG--VFSVLYTVP
                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                            4.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13:
                                                                                                                                                                                                                                                                                                             40;
                                                                                           ---KGSW----
                                                                                                                                                                                    ----QVDGDLLSGVCYVGLSSVDALRGFVLAPLFVYLF-----
                                                                                                                                                                                                                                                                                                               ; Score 96; DB; Pred. No. 0.22
                                                                                                                                                                                                                                                                                                                                            DB 4;
                                                                                                                                                                                                                                                                                                             79;
                                                                                                                                                                                                                                                                                                                                          Length 572;
                                                                                                                                                                                                                                                                                                                Indels
                                                                                           -NSDITLLTISFIFPLI 236
                                                                                                                                                                                                                                                                                                               84;
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Gaps

Search completed: June 9 2003, 07:10:42

537

479

432